Att'y Dkt. No.: US-1420

U.S. App. No: 09/897,988

## IN THE CLAIMS:

Kindly rewrite Claims 1-10 as follows, in accordance with 37 C.F.R. § 1.121:

1. (currently amended) A method [[,]] comprising:

culturing a microorganism in a medium to produce and cause accumulation of a target substance in the medium; and

collecting the target substance,

wherein the microorganism is constructed from a parent strain of the microorganism belonging to the genus Escherichia or Coryneform bacteria and having a respiratory chain pathway of high energy efficiency and a respiratory chain pathway of low energy efficiency as respiratory chain pathways, and

the microorganism is a mutant strain or a genetic recombinant strain having either one or both of the following characteristics:

- (A) the activity of an enzyme of the respiratory chain pathway of high energy efficiency selected from the group consisting of SoxM type oxidase, bc1 complex, cytochrome bo-type oxidase, and NDH-1 is enhanced,
- (B) the activity of an enzyme of the respiratory chain pathway of low energy efficiency selected from the group consisting of cytochrome bd type oxidase and NDH-II is deficient.

wherein the target substance is selected from the group consisting of an L-amino acid and a nucleic acid.

2. (previously presented) The method according to Claim 1, wherein the activity of an enzyme of the respiratory chain pathway of high energy efficiency is enhanced by a method selected from the group consisting of

increasing a copy number of a gene coding for said enzyme; and modifying an expression regulatory sequence of said gene.

- 3. (previously presented) The method according to Claim 1, wherein the activity of an enzyme of the respiratory chain pathway of low energy efficiency is made deficient by disruption of a gene coding for said enzyme.
- 4. (cancelled)
- 5. (cancelled)

Att'y Dkt. No.: US-1420

U.S. App. No: 09/897,988

- 6. (previously presented) The method according to Claim 1, wherein said microorganism comprises enhanced SoxM type oxidase activity and deficient NDH-II activity.
- 7. (previously presented) The method according to Claim 1, wherein an enzyme of the respiratory chain pathway of high energy efficiency is cytochrome bo type oxidase.
- 8. (cancelled)
- 9. (cancelled)
- 10. (currently amended) The method according to Claim <u>§1</u>, wherein the microorganism is a bacterium belonging to the genus Escherichia.